

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/480,494DATE: 02/05/97  
TIME: 16:03:36

INPUT SET: S15332.raw

This Raw Listing contains the General  
Information Section and those Sequences  
containing ERRORS.

Does Not Comply  
Corrected Diskette Needed

1  
2  
3  
4  
5  
6  
7  
8  
9  
10  
11  
12  
13  
14  
15  
16  
17  
18  
19  
20  
21  
22  
23  
24  
25  
26  
27  
28  
29  
30  
31  
32  
33  
34  
35  
36  
37  
38  
39  
40

SEQUENCE LISTING

(1) General Information:

(i) APPLICANT: Roeske, Roger W.

(ii) TITLE OF INVENTION: LHRH Antagonist Peptides

(iii) NUMBER OF SEQUENCES: 1

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: LAHIVE & COCKFIELD  
(B) STREET: 60 State Street, Suite 510  
(C) CITY: Boston  
(D) STATE: Massachusetts  
(E) COUNTRY: USA  
(F) ZIP: 02109-1875

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk  
(B) COMPUTER: IBM PC compatible  
(C) OPERATING SYSTEM: PC-DOS/MS-DOS  
(D) SOFTWARE: PatentIn Release #1.0, Version #1.25

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: 08/480,494  
(B) FILING DATE: 07-JUN-1995  
(C) CLASSIFICATION:

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: DeConti, Giulio A.  
(B) REGISTRATION NUMBER: 31,503  
(C) REFERENCE/DOCKET NUMBER: PPI-007

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: (617)227-7400  
(B) TELEFAX: (617)227-5941

ERRORED SEQUENCES FOLLOW:

RAW SEQUENCE LISTING  
PATENT APPLICATION *US/08/480,494*

DATE: 02/05/97

TIME: 16:03:37

INPUT SET: S15332.raw

41 (2) INFORMATION FOR SEQ ID NO:1:

42

43 (i) SEQUENCE CHARACTERISTICS:

--&gt; 44 (A) LENGTH: 10 amino acids

45 (B) TYPE: amino acid

46 (D) TOPOLOGY: linear

47

48 (ii) MOLECULE TYPE: peptide

49

50 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

51

--&gt; 52 Glu-His-Trp-Ser-Tyr-Gly-Leu-Arg-Pro-Gly

53

1

5

10

--> 54 -(PAGE|34)-

55

56

57

58

*delete  
also**please  
delete dashes —  
dashes are not  
permitted*

SEQUENCE VERIFICATION REPORT  
PATENT APPLICATION US/08/480,494

DATE: 02/05/97

TIME: 16:03:39

INPUT SET: S15332.raw

Line	Error	Original Text
44	Entered (10) and Calc. Seq. Length (0) differ	(A) LENGTH: 10 amino acids
52	Wrong Amino Acid Designator	Glu-His-Trp-Ser-Tyr-Gly-Leu-Arg-Pro-Gly
54	Wrong Amino Acid Designator	-(PAGE 34)-

## Notice of Availability

### Applicant Aid for Biotechnology Computer Readable Form (CRF) Sequence Listings Submissions

The Patent and Trademark Office (PTO) has developed a computer program, called Checker, that will aid applicants in identifying and correcting errors prior to making submissions for compliance with the Requirements for Patent Applications Containing Nucleotide Sequence and/or Amino Acid Sequence Disclosures (sequence rules: 37 CFR 1.821 through 1.825). (Final rules were published in the *Federal Register* (55 FR 18230) on May 1, 1990, and in the *PTO Official Gazette* (1114 Off.Gaz.PatOffice 29) on May 15, 1990.)

Checker is a DOS-based software program that is intended to assist users in determining whether errors may be present in the sequence listings, and is not intended to guarantee that the submission is error-free.

The most current version of the software will be available via computer downloading (details below). Copies on diskette are also available. Updated software versions will not be automatically mailed out; any updates will be announced in the *PTO Official Gazette*.

The software can be accessed/requested in the following locations:

- 1) Dial-up access to the Patent and Trademark Office Bulletin Board System.  
Phone number: 703-305-8950  
Cost: Free-of-charge
- 2) Dial-up access through the Internet. FTP site: ftp.uspto.gov  
Login as "anonymous". Software is in directory /pub/checker  
Cost: Free-of-charge
- 3) For diskette copies, telephone requests to 703-306-2600.  
Cost: \$25.00

For Further Information Contact: Meredith Beckhardt at 703-308-4212